```
                      GenCore version 6.3
             Copyright (c) 1993 - 2009  Biocceleration Ltd.



OM nucleic - nucleic search, using sw model

Run on:          June 26, 2009, 03:19:14 ; Search time 752 Seconds
                                            (without alignments)
                                      14891.617 Million cell
updates/sec

Title:           US-09-300-482-298
Perfect score:   388
Sequence:        1
ggagaaagaaagaaaagatg..........ttaaagcctgggacccccgt 388

Scoring table:   IDENTITY_NUC
                 Gapop 10.0 , Gapext 1.0

Searched:        41078765 seqs, 14427166270 residues

Total number of hits satisfying chosen parameters:      82157530

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

Database :       Published_Applications_NA_Main:*
            1:
/ABSS/Data/CRF/ptodata/1/pubpna/US07_PUBCOMB.seq:*
            2:
/ABSS/Data/CRF/ptodata/1/pubpna/US08_PUBCOMB.seq:*
            3:
/ABSS/Data/CRF/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
            4:
/ABSS/Data/CRF/ptodata/1/pubpna/US09B_PUBCOMB.seq:*
            5:
/ABSS/Data/CRF/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
            6:
/ABSS/Data/CRF/ptodata/1/pubpna/US09D_PUBCOMB.seq:*
            7:
/ABSS/Data/CRF/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
```